- 1. An antibody-based fusion protein comprising an immunoglobulin (Ig) chain linked to a non-Ig protein via a junction point, wherein said antibody-based fusion protein comprises an amino acid alteration within 10 amino acids from said junction point, in said Ig chain or said non-Ig protein, and wherein said antibody-based fusion protein has a longer circulating half-life *in vivo* than a corresponding antibody-based fusion protein without said amino acid alteration.
- 2. The fusion protein of claim 1 wherein the amino acid alteration increases the hydrophobicity of said antibody-based fusion protein.
- 3. The fusion protein of claim 1 or 2 wherein said Ig chain is N-terminal to said nonIg protein.
- 4. The fusion protein of claim 1,2 or 3 wherein said alteration changes the C-terminal amino acid of the Ig chain.
- 5. The fusion protein of claim 1, wherein said non-Ig protein is a secreted protein.
- 6. The fusion protein of claim 5, wherein said non-Ig protein is a mature form of said secreted protein.
- 7. The fusion protein of claim 1, wherein the Ig chain comprises part of an Ig heavy chain.
- 8. The antibody-based fusion protein of claim 7 wherein said Ig chain comprises at least the CH2 domain of an IgG2 or an IgG4 constant region.
- 9. The antibody-based fusion protein of claim 7, wherein said Ig chain comprises at least a portion of an IgG1 constant region having a mutation or a deletion at one or more amino acids selected from the group consisting of Leu₂₃₄, Leu₂₃₅, Gly₂₃₆, Gly₂₃₇, Ash₂₉₇, and Pro₃₃₁.
- 10. The antibody-based fusion protein of claim 7, wherein said Ig chain comprises at least a portion of an IgG3 constant region having a mutation or a deletion at one or more amino acids selected from the group consisting of Leu₂₈₁, Leu₂₈₂, Gly₂₈₃, Gly₂₈₄, Asn₃₄₄, and Pro₃₇₈.

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- 11. The antibody-based fusion protein of claim 7, wherein said Ig chain has binding affinity for an immunoglobulin protection receptor.
- 12. The antibody-based fusion protein of claim 7, wherein said Ig chain has substantially reduced binding affinity for a Fc receptor selected from the group consisting of FcγRI, FcγRII and FcγRIII.
- 13. The antibody-based fusion protein of claim 7, wherein said non-Ig protein is selected from the group consisting of a cytokine, a ligand-binding protein, and a protein toxin.
- 14. The antibody-based fusion protein of claim 13, wherein said cytokine is selected from the group consisting of a tumor necrosis factor, an interleukin, and a lymphokine.
- 15. The antibody-based fusion protein of <u>claim</u> 14, wherein said tumor necrosis factor is tumor necrosis factor alpha.
- 16. The antibody-based fusion protein of claim 14, wherein said interleukin is interleukin-2.
- 17. The antibody-based fusion protein of <u>claim</u> 14, wherein said lymphokine is a lymphotoxin or a colony stimulating factor.
- 18. The antibody-based fusion protein of claim 11, wherein said colony stimulating factor is a granulocyte-macrophage colony stimulating factor.
- 19. The antibody-based fusion protein of claim 13, wherein said ligand-binding protein is selected from the group consisting of CD4, CTLA-4, TNF receptor, and an interleukin receptor.

A method for increasing the circulating half-life of an antibody-based fusion protein having an Ig chain linked to a non-Ig protein via a junction point, the method comprising the step of substituting, deleting, inserting, or otherwise altering an amino acid at or near said junction point.

The method of <u>claim</u> 20, wherein said fusion protein comprises a portion of a heavy chain.

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∫21.

- 22. The method of claim 21, wherein said fusion protein comprises at least the CH2 domain of an IgG2 or an IgG4 constant region.
- 23. The method of claim 20, 21, or 22, wherein said fusion protein comprises a heavy chain moiety having a mutation that affects interaction with an Fc protection receptor.
- 24. The fusion protein of claim I comprising a linker between said Ig chain and said non-Ig protein.
- 25. The fusion protein of claim 4, 5, 6, or 7, wherein said alteration is a substitution of one or more amino acids.
- 26. An antibody-based fusion protein comprising
 - (a) a first polypeptide comprising an Ig chain, and,
 - b) a second polypeptide comprising a non-Ig protein,

wherein said first polypeptide is joined to said second polypeptide to produce a junction region having at least one mutation, and

wherein said fusion protein has a longer circulating half life than a fusion protein having a junction region without said mutation.

- 27. The fusion protein of claim 26, wherein said mutation is in the C-terminal portion of said first polypeptide
- 28. The fusion protein of claim 26, wherein said mutation is in the N-terminal portion of said second polypeptide.
- 29. The fusion protein of claim 26 comprising a first mutation in the C-terminal portion of said first polypeptide and a second mutation in the N-terminal portion of said second polypeptide.
- 30. The fusion protein of claim 27 or 29 wherein said C-terminal portion comprises between 1 and 100 C-terminal amino acids of said first polypeptide.
- 31. The fusion protein of claim 30 wherein said C-terminal portion comprises between 1 and 10 C-terminal amino acids of said first polypeptide.
- 32. The fusion protein of claim 28 or 29 wherein said N-terminal portion comprises between 1 and 100 N-terminal amino acids of said second polypeptide.

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- 33. The fusion protein of claim 32 wherein said N-terminal portion comprises between 1 and 10 N-terminal amino acids of said second polypeptide.
- 34. The fusion protein of claim 26 wherein said Ig is IgG1.
- 35. The fusion protein of claim 26 wherein said mutation is selected from the group consisting of point mutations, deletions, insertions, and rearrangements.
- 36. The fusion protein of claim 34 wherein the C-terminal residue of said first polypeptide is mutated to be an amino acid with a non-ionizable side chain.
- 37. The fusion protein of claim 36 wherein said C-terminal residue is a non-lysine amino acid.
- 38. The fusion protein of claim 26 wherein said junction region consists of the C-terminal region of said first polypeptide and the N-terminal region of said second polypeptide, and wherein said mutation is present in one of said C-terminal and N-terminal regions.

The fusion protein of claim 26 wherein said junction region comprises a spacer or linker peptide.

- The fusion protein of claim 39 wherein said mutation consists of the presence of a spacer or linker peptide between said first and said second polypeptides.
- 41. The fusion protein of claim 26 wherein said mutation is in a region that does not interact with FcR or FcRp.
- 42. A method for identifying a mutation that increases the circulating half life of an antibody-based fusion protein having an Ig moiety and a non-Ig moiety comprising the steps of:
 - a) introducing a mutation in the region spanning the junction between the
 Ig moiety and the non-Ig moiety;
 - b) comparing the serum half-lives of the antibody-based fusion protein with and without a mutation; and,
 - c) selecting a mutation that increases the serum half-life of the antibodybased fusion protein.

- 43. An antibody-based fusion protein comprising a mutation identified according to the method of claim 42.
- 44. A method for treating a disease comprising the step of administering to a patient an antibody-based fusion protein of claim 26.
- 45. The fusion protein of claim 1, 26, or 36, having a hydrophobic or non-polar amino acid introduced via addition or substitution at or near said junction.
- 46. The fusion protein of claim 45, wherein said amino acid is selected from the group consisting of Leu, Ala, Tri, and Gly.
- 47. The fusion protein of claim 46, wherein said amino acid is Ala.

